

SEQUENCE LISTING

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ABE, Kunitake
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<120> NOVEL PROMOTER

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<140> 10/509,565

<141> 2004-09-29

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<170> PatentIn version 3.3

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Leu His Leu Arg Leu Lys Gly Ser Arg His Asp Phe His Val Asp Leu
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370

375

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Pro Val Lys Glu Tyr Lys Tyr Pro Glu Lys Leu Pro Gly Glu Leu Tyr
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Asp Ala Asn Thr Gln Cys Lys Trp Gln Phe Gly Glu Lys Ala Lys Leu
 515 520 525

Cys Met Leu Asp Phe Lys Lys Asp Ile Cys Lys Ala Leu Trp Cys His
 530 535 540

Arg Ile Gly Arg Lys Cys Glu Thr Lys Phe Met Pro Ala Ala Glu Gly
 545 550 555 560

Thr Ile Cys Gly His Asp Met Trp Cys Arg Gly Gly Gln Cys Val Lys
 565 570 575

Tyr Gly Asp Glu Gly Pro Lys Pro Thr His Gly His Trp Ser Asp Trp
 580 585 590

Ser Ser Trp Ser Pro Cys Ser Arg Thr Cys Gly Gly Gly Val Ser His
 9/21

595

600

605

Arg Ser Arg Leu Cys Thr Asn Pro Lys Pro Ser His Gly Gly Lys Phe
 610 615 620

Cys Glu Gly Ser Thr Arg Thr Leu Lys Leu Cys Asn Ser Gln Lys Cys
 625 630 635 640

Pro Arg Asp Ser Val Asp Phe Arg Ala Ala Gln Cys Ala Glu His Asn
 645 650 655

Ser Arg Arg Phe Arg Gly Arg His Tyr Lys Trp Lys Pro Tyr Thr Gln
 660 665 670

Val Glu Asp Gln Asp Leu Cys Lys Leu Tyr Cys Ile Ala Glu Gly Phe
 675 680 685

Asp Phe Phe Phe Ser Leu Ser Asn Lys Val Lys Asp Gly Thr Pro Cys
 690 695 700

Ser Glu Asp Ser Arg Asn Val Cys Ile Asp Gly Ile Cys Glu Arg Val
 705 710 715 720

Gly Cys Asp Asn Val Leu Gly Ser Asp Ala Val Glu Asp Val Cys Gly
 725 730 735

Val Cys Asn Gly Asn Asn Ser Ala Cys Thr Ile His Arg Gly Leu Tyr
 740 745 750

Thr Lys His His His Thr Asn Gln Tyr Tyr His Met Val Thr Ile Pro
 755 760 765

Ser Gly Ala Arg Ser Ile Arg Ile Tyr Glu Met Asn Val Ser Thr Ser
 770 775 780

Tyr Ile Ser Val Arg Asn Ala Leu Arg Arg Tyr Tyr Leu Asn Gly His
 785 790 795 800

Trp Thr Val Asp Trp Pro Gly Arg Tyr Lys Phe Ser Gly Thr Thr Phe
 805 810 815

Asp Tyr Arg Arg Ser Tyr Asn Glu Pro Glu Asn Leu Ile Ala Thr Gly

820

825

830

Pro Thr Asn Glu Thr Leu Ile Val Glu Leu Leu Phe Gln Gly Arg Asn
835 840 845

Pro Gly Val Ala Trp Glu Tyr Ser Met Pro Arg Leu Gly Thr Glu Lys
850 855 860

Gln Pro Pro Ala Gln Pro Ser Tyr Thr Trp Ala Ile Val Arg Ser Glu
865 870 875 880

Cys Ser Val Ser Cys Gly Gly Gly Gln Met Thr Val Arg Glu Gly Cys
885 890 895

Tyr Arg Asp Leu Lys Phe Gln Val Asn Met Ser Phe Cys Asn Pro Lys
900 905 910

Thr Arg Pro Val Thr Gly Leu Val Pro Cys Lys Val Ser Ala Cys Pro
915 920 925

Pro Ser Trp Ser Val Gly Asn Trp Ser Ala Cys Ser Arg Thr Cys Gly
930 935 940

Gly Gly Ala Gln Ser Arg Pro Val Gln Cys Thr Arg Arg Val His Tyr
945 950 955 960

Asp Ser Glu Pro Val Pro Ala Ser Leu Cys Pro Gln Pro Ala Pro Ser
965 970 975

Ser Arg Gln Ala Cys Asn Ser Gln Ser Cys Pro Pro Ala Trp Ser Ala
980 985 990

Gly Pro Trp Ala Glu Cys Ser His Thr Cys Gly Lys Gly Trp Arg Lys
995 1000 1005

Arg Ala Val Ala Cys Lys Ser Thr Asn Pro Ser Ala Arg Ala Gln
1010 1015 1020

Leu Leu Pro Asp Ala Val Cys Thr Ser Glu Pro Lys Pro Arg Met
1025 1030 1035

His Glu Ala Cys Leu Leu Gln Arg Cys His Lys Pro Lys Lys Leu

1040		1045		1050
Gln Trp Leu Val Ser Ala Trp Ser Gln Cys Ser Val Thr Cys Glu				
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Arg Gly Thr Gln Lys Arg Phe Leu Lys Cys Ala Glu Lys Tyr Val				
1070		1075		1080
Ser Gly Lys Tyr Arg Glu Leu Ala Ser Lys Lys Cys Ser His Leu				
1085		1090		1095
Pro Lys Pro Ser Leu Glu Leu Glu Arg Ala Cys Ala Pro Leu Pro				
1100		1105		1110
Cys Pro Arg His Pro Pro Phe Ala Ala Ala Gly Pro Ser Arg Gly				
1115		1120		1125
Ser Trp Phe Ala Ser Pro Trp Ser Gln Cys Thr Ala Ser Cys Gly				
1130		1135		1140
Gly Gly Val Gln Thr Arg Ser Val Gln Cys Leu Ala Gly Gly Arg				
1145		1150		1155
Pro Ala Ser Gly Cys Leu Leu His Gln Lys Pro Ser Ala Ser Leu				
1160		1165		1170
Ala Cys Asn Thr His Phe Cys Pro Ile Ala Glu Lys Lys Asp Ala				
1175		1180		1185
Phe Cys Lys Asp Tyr Phe His Trp Cys Tyr Leu Val Pro Gln His				
1190		1195		1200
Gly Met Cys Ser His Lys Phe Tyr Gly Lys Gln Cys Cys Lys Thr				
1205		1210		1215
Cys Ser Lys Ser Asn Leu				
1220				

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 <210> 4
 <211> 50
 <212> DNA
 <213> Artificial Sequence

 <220>
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 <400> 4
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 <210> 5
 <211> 34
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> an artificially synthesized primer sequence

 <400> 5
 ggactagtct agaagctggg taccagctgc tagc 34

 <210> 6
 <211> 29
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> an artificially synthesized primer sequence

 <400> 6
 ggactagtgt cgaccggtca tggctgcgc 29

 <210> 7
 <211> 38
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> an artificially synthesized primer sequence

 <400> 7
 ggactagtgc catgggaccc gcagcggcag cgcttggg 38

 <210> 8

<211> 40
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<213> Artificial Sequence

<220>
<223> an artificially synthesized primer sequence

<400> 8
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<211> 41
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<213> Artificial Sequence

<220>
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<400> 9
ggactagtac catgaagccc cgcgcgcgcg gatggcgggg c 41

<210> 10
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<213> Homo sapiens

<400> 10
ccctgtggtc aacctcgtag gcagagacca 30

<210> 11
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<212> PRT
<213> Artificial Sequence

<220>
<223> peptide expressed from polynucleotide comprising NotI recognition
sequence and FLAG tag

<400> 11

Ala Ala Ala Asp Tyr Lys Asp Asp Asp Asp Lys
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<210> 12
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<400> 12

His Glu Ser Gly His
1 5

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 <220>
 <223> an artificially synthesized primer sequence

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 <213> Homo sapiens

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 gaggaaggaa caaaattaca gactaaacag aatctttgta gggcacaaac acatccaatt 180

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 <213> Homo sapiens

<400> 18
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<210> 19
 <211> 29
 <212> DNA
 <213> Homo sapiens

<400> 19
 ccaggagcgc tccgaggggc taggggcca 29

<210> 20
 <211> 43
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> an artificially synthesized primer sequence

<400> 20
 aagagctctg ctagctgaga cggagtctcg ctctgtcgcc cag 43

<210> 21
 <211> 43
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> an artificially synthesized primer sequence

<400> 21
 taaagcttag atctccagga gcgctccgag gggctagggg cca 43

<210> 22
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 <212> DNA
 <213> Artificial Sequence

<220>
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 <210> 26
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 <400> 26
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 <210> 27
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 <400> 28
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 <400> 29
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<400> 30
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19

<210> 31
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<400> 31
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21

<210> 32
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15

<210> 33
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21